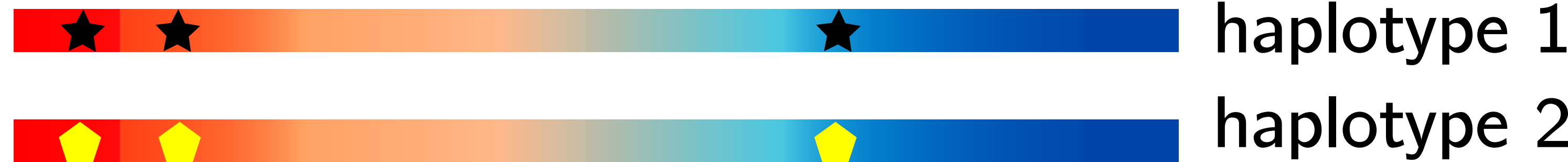


100kb region from the genome



1.000.000.000 reads → 100.000 contigs

